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54) Title: METHOD OF SYNTHESIZING  57) Abstract  A general stochastic method for syntheroperties. The use of identification tags on	esizing random o	igomers can be used to synthesize compou	nds to screen for desire
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## METHOD OF SYNTHESIZING DIVERSE COLLECTIONS OF OLIGOMERS

## FIELD OF THE INVENTION

The present invention relates generally to stochastic methods for synthesizing random oligomers, with particular emphasis on particle-based synthesis methods. The invention also relates to the use of identification tags on the particles to facilitate identification of the oligomer sequence synthesized. Yet another aspect of the invention relates to the use of tagged oligomer libraries in receptor-binding studies.

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## BACKGROUND OF THE INVENTION

The relationship between structure and activity of molecules is a fundamental issue in the study of biological systems. Structure-activity relationships are important in understanding, for example, the function of enzymes, the ways in which cells communicate with each other, and cellular control and feedback systems. Certain macromolecules are known to interact and bind to other molecules having a very specific three-dimensional spatial and electronic distribution. Any large molecule having such specificity can be considered a receptor, whether the molecule is an enzyme catalyzing hydrolysis of a metabolic intermediate, a cell-surface protein mediating membrane transport of ions, a glycoprotein serving to identify a particular cell to its neighbors, an IgG-class antibody circulating in the plasma, an oligonucleotide sequence of DNA in the genome, or the like. The various molecules that receptors selectively bind are known as ligands.

Many assays are available for measuring the binding affinity of known receptors and ligands, but the information that can be gained from such experiments is often limited by the number and type of available ligands. Novel ligands are sometimes discovered by chance or by application of new techniques for the elucidation of molecular structure, including x-ray crystallographic analysis and recombinant genetic techniques for proteins.

Small peptides are an exemplary system for exploring the relationship between structure and function in biology. A peptide is a polymer composed of amino acid monomers. When the twenty naturally occurring amino acids are condensed into polymeric molecules, the resulting polymers form a wide variety of three-dimensional configurations, each resulting from a particular amino acid sequence and solvent condition. The number of possible pentapeptides of the 20